SEQUENCE LISTING

<110> Barber, Elizabeth K <120> Gene Expression Control Element DNA <130> 896034605001 RECEIVED <140> US 09/966,264 AUG 2 2 2002 <141> 2001-09-28 <150> US 60/237,079 TECH CENTER 1600/2900 <151> 2000-09-30 <160> 33 <170> PatentIn version 3.1 <210> 1 <211> 137 <212> DNA <213> human <400> 1 at tat aaa gga aaa aga aaa taa cgc aat gga caa gtg gtg aag ctg 47 Tyr Lys Gly Lys Arg Lys Arg Asn Gly Gln Val Val Lys Leu 1 10 tga act cag gtg tgc aca att atc agg aac acc cca aaa cca aag tga 95 Thr Gln Val Cys Thr Ile Ile Arg Asn Thr Pro Lys Pro Lys 15 20 ggt aga aat agc atg aga agc cgt gtt tga tgt taa tta att 137 Gly Arg Asn Ser Met Arg Ser Arg Val Cys Leu Ile 40 <210> 2 <211> 996 <212> DNA <213> human <400> 2 gtg gtt tga ttg ata gta aaa aaa atg ttc gtt aat aca agt aga gag 48 Val Val Leu Ile Val Lys Lys Met Phe Val Asn Thr Ser Arg Glu 10 15 taa gta atc aat caa tca ctc ata gcc aag gtg gaa aag atg tat ccc 96 Val Ile Asn Gln Ser Leu Ile Ala Lys Val Glu Lys Met Tyr Pro

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Gln Arg Gly Leu Asp Ser Lys Ser Leu Gln Glu Ile Asn Leu Tyr Phe
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Asn Ile Lys Leu Ile Leu Thr Asn Asn Val Lys Trp Leu His Lys 50 55 60

Lys	Gly	Phe	Ala	Ser 65	Ser	Trp	Lys	Leu	Val 70	Lys	Asn	Gln	Thr	Leu 75
Leu	Сув	Thr	Pro	Ser 80	Met	Gln	Leu	Leu	Сув 85	Cys	Leu	His	Pro	Glu 90
Met	Gly	Asn	Asp	Phe 95	Pro	Asn	Gly	Lys	Glu 100	Thr	Glu	Arg	Cys	Tyr 105
Leu	Ser	Ala	Pro	Phe 110	Val	Lys	Ser	Val	Phe 115	Leu	Ser	Leu	Cys	Phe 120
Pro	Gly	His	Asn	Val 125	Gly	Ser	Leu	Phe	His 130	Met	Ala	Asp	Asp	Leu 135
Gly	Arg	Ala	Met	Glu 140	Ser	Leu	Val	Ser	Val 145	Met	Thr	Asp	Glu	Glu 150
Gly	Ala	Glu	Lys	Met 155	Phe	Tyr	Asn	Ser	Arg 160	Phe	Pro	His	Gly	Phe 165
Tyr	Asn	Ile	His	Thr 170	Thr	Lys	Arg	Ile	Arg 175	Gln	Lys	Glu	Phe	Thr 180
Arg	Asn	Lys	Ser	Ile 185	Phe	Leu	Arg	Arg	Val 190	Val	Val	Leu	Tyr	Cys 195
Arg	Phe	Gln	Lys	Phe 200	Leu	Ser	Leu	Leu	Leu 205	Phe	Cys	Lys	G1n	Trp 210
Gln	Val	Leu	His	Val 215	Tyr	Ala	Ile	Val	Gln 220	Lys	Ser	Tyr	Lys	Lys 225
Thr	Thr	Cys	Lys	Ile 230	Leu	Ile	Ala	Lys	Lys 235	Leu	Ala	Ile	Ser	Leu 240
Tyr	Gly	Thr	His	Phe 245	Gly	Leu	Phe	Lys	Asn 250	Leu	Lys	Gln	Leu	Lys 255
Arg	Lys	Asn	Tyr	Lys 260	Gly	Lys	Arg	Lys	Lys 265	Arg	Asn	Gly	Gln	Val 270
Val	T	Leu	Ara	Thr	Gln	Val	Cys	Thr	Ile	Ile	Arg	Asn	Thr	Pro
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Lys Leu Ile

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ATTGCG	TAT	TTTCTTTTTC	CTTTATAATT	CTTTCTTTTT	CCTTCATAAT	200
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Asn	Met	Lys	Lys	Glu 35	Gln	Asn	Lys	Cys	Phe 40	Thr	Thr	Pro	Asp	Ser 45
Arg	Met	Val	Phe	Ile 50	Ile	Phe	Ile	Gln	Gln 55	Arg	Gly	Leu	Asp	Ser 60
Lys	Ser	Leu	Gln	Glu 65	Ile	Asn	Leu	Tyr	Phe 70	Cys	Glu	Gly	Phe	Tyr 75
Thr	Ser	Met	Gln	Leu 80	Tyr	Lys	Lys	Val	Ile 85	Arg	Lys	Leu	His	Lys 90
Ile	Thr	Gln	Trp	Thr	Arg	Thr	Pro	Gln	Asn	Gln	Ser	Glu	Val	Glu
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Ile Ala